FIGURE 1

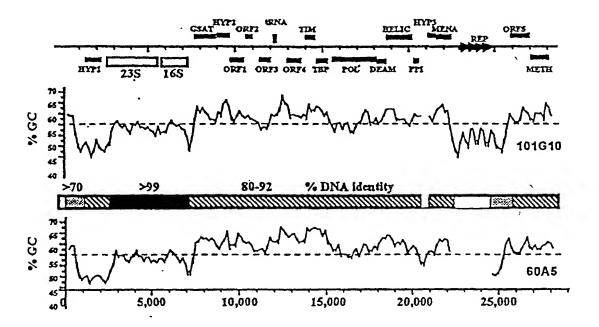


Figure 2

vo.	<u>Gene</u>	<u>Strain</u>		TATA BOX			Coding Start		TATA to Start (bp)	
81 82	Hypoth 03	A B					GCGGCGCATG CCGGCGCGTG			25
83 84	Hypoth 02	A B	GGAAACTTTG	ATTATA	CGGG	CGTACATTCC	CGGGGCCCAT CGGGGCCCAT	G~~~~~	~~~~~~~	26
85 86	ORF 02	A B	ACGGCAAGGT	AATAAT	AGCC	TGCCGTCCGT	AACGGCCGTA ACCTGCCGTA	TG	~~~~~~~	27
87 88		A B	CATGGAACTA	GATAAT	AACC	GGTCCCGCGG	ATCCCATGCA GTACAATGCA	TG~~~~~~	~~~~~~~	27
90	regal to specific to g	A B	AGCACGACAA	GTTATA	GCAG	GGTACAAAGG	GTGCGCGCGC AGCAGCGCAC GCCTGCTGCC	ATG	~~~~~~~	28
92		A B	ATCCGGCCTC	ATTAAA	TTAC	GGGGGTACA	ACCTGCTGCC GCGGCTGCGC	GTG~~~~~	~~~~~~	28
94		A B	ACTTCATACA	CATAAA	TCCC	GCCTGAACGG	TCGTCCGCGC CACCATGGCC	ATG	~~~~~~~	29
96	deaminase	A B A	CCGCATATAC	CATAAT	ATGC	CGGGCGGGG	CAGGCTGCCC CAGGGCCGCG	.GTG~~~~~	~~~~~~~	29
98	ORF 06	B A	GGGTAGAAAC	CATAAA	ACAA	CAGGCCGCGG	CAGGGCG.CG GCGCGTATCA	CGTG~~~~~	~~~~~~~	29
100	tRNA-tyr	B A	GCGATAGTTA	TTTAAA	ACTA	GGATGCCGAT	GCGCGGACCA CACGGATCGT	CCCA~~~~~	~~~~~~~	29
103	± TBP	B A	CCGGGCCCCG	GTTAAA	ATAG	CG.CACGGGC	CACCCGTCGT GGATCCTGAC	CAATG~~~~	~~~~~~	30
104	TIM	B A	GCGTCGATAG	AATAAA	TACG	CGCAGGGGGC	GGCACCGGAT CCCGTGGCGC GCGGTGC	GATCGCCCGT	G~~~~~	36
107		B A B	ATTTCAACTA	CATAAA	TGCC	TAGTTACGCA	GAAATAGCAA GAAATATCAA	ACGACGTACT	TCGACTAATG	45
109	ORF 01	A B					TGTA //G TACA //G			52
111	Methylase	В	CTACAAAGAT	TTTAAG	ACGG	CGCGGGTGCC	GCCG.//G GCGG.//T	GGCACGGGGG	CCTATCTTG	104
113	16S RNA						CCGATCCGAT GCGATCCGAT			220
·	Archaeal promoter consensus			YTTAWA	l.					

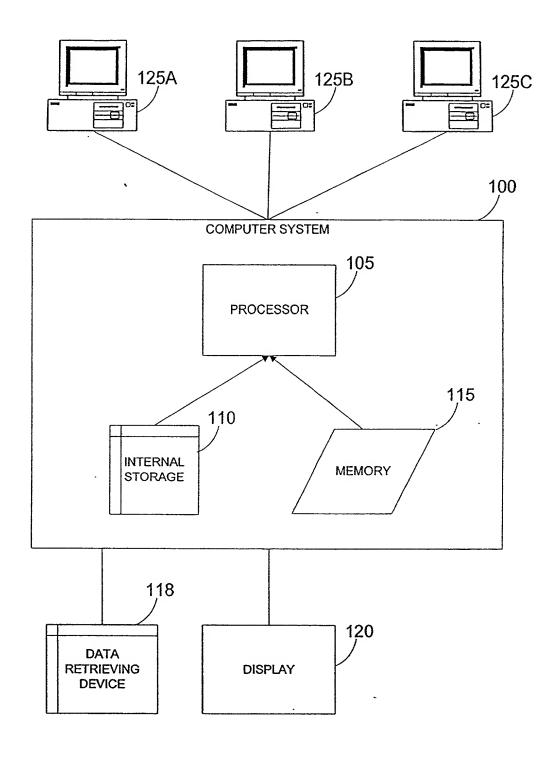


FIGURE 3

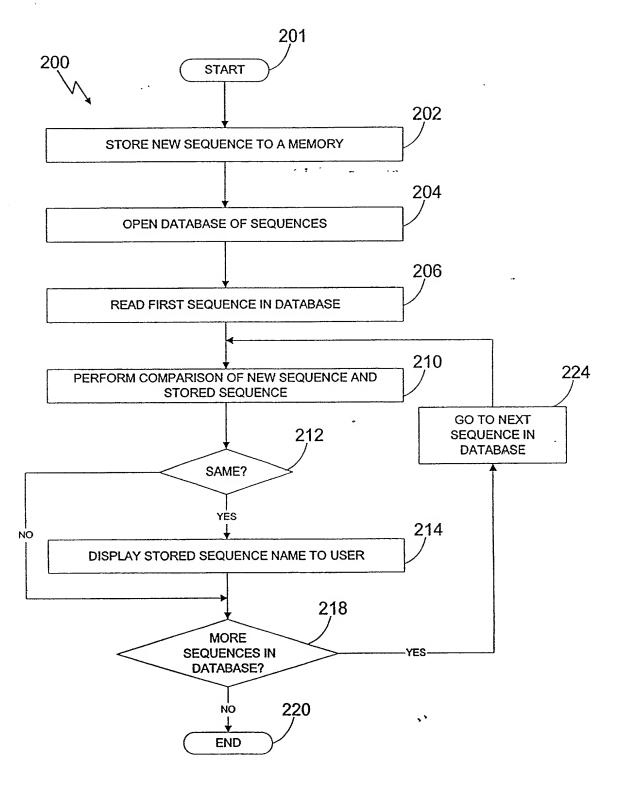


FIGURE 4

